



OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/957,458A

DATE: 03/24/2003  
TIME: 13:43:57

Input Set : A:\09957458.ST25.txt  
Output Set: N:\CRF4\03242003\I957458A.raw

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3 <110> APPLICANT: chen, una
5 <120> TITLE OF INVENTION: Method for growing stem cells
7 <130> FILE REFERENCE: P66567US0
9 <140> CURRENT APPLICATION NUMBER: US 09/957,458A
10 <141> CURRENT FILING DATE: 2001-09-21
12 <150> PRIOR APPLICATION NUMBER: PCT/EP00/08247
13 <151> PRIOR FILING DATE: 2000-08-24
15 <150> PRIOR APPLICATION NUMBER: EP 99116533
16 <151> PRIOR FILING DATE: 1999-08-24
18 <160> NUMBER OF SEQ ID NOS: 10
20 <170> SOFTWARE: PatentIn version 3.2
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23 <211> LENGTH: 7969
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62 <221> NAME/KEY: misc_feature

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ppr 1,6  
Does Not Comply  
Corrected Diskette Needed

insufficient response give source of  
genetic material.  
See item 11  
on Error Summary  
Sheet

Please correct  
any similar  
errors in  
subsequent  
sequences.

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63 <222> LOCATION: (762)..(762) /  
64 <223> OTHER INFORMATION: n is a, c, g, or t  
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74 <223> OTHER INFORMATION: n is a, c, g, or t  
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77 <221> NAME/KEY: misc\_feature  
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Input Set : A:\09957458.ST25.txt  
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124 <223> OTHER INFORMATION: n is a, c, g, or t  
126 <220> FEATURE:  
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159 <223> OTHER INFORMATION: n is a, c, g, or t  
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182 <221> NAME/KEY: misc\_feature  
183 <222> LOCATION: (1040)..(1041) /  
184 <223> OTHER INFORMATION: n is a, c, g, or t

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Input Set : A:\09957458.ST25.txt  
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186 <220> FEATURE:  
187 <221> NAME/KEY: misc\_feature  
188 <222> LOCATION: (1047)..(1047) /  
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221 <220> FEATURE:  
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224 <223> OTHER INFORMATION: n is a, c, g, or t  
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229 <223> OTHER INFORMATION: n is a, c, g, or t  
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233 <222> LOCATION: (1120)..(1120) /  
234 <223> OTHER INFORMATION: n is a, c, g, or t  
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237 <221> NAME/KEY: misc\_feature  
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242 <221> NAME/KEY: misc\_feature  
243 <222> LOCATION: (1126)..(1126) /  
244 <223> OTHER INFORMATION: n is a, c, g, or t  
246 <220> FEATURE:

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247 <221> NAME/KEY: misc_feature
248 <222> LOCATION: (1133)..(1133)
249 <223> OTHER INFORMATION: n is a, c, / g, or t
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262 <221> NAME/KEY: misc_feature
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276 <400> SEQUENCE: 1

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|      | 279 | tgtgacggtc cttacaatga aatgcancgt gtttatctt ttcctgtatgg cagggttac     | 120  |
|      | 281 | aggtaagggg ctcccaagtc ccaaacttga gggccataa actctgtgac agtggcaatc     | 180  |
|      | 283 | actttgcott tcttctaca ggggtgaatt cggcttcac agagcattca ccgctgaccc      | 240  |
|      | 285 | ctcaccgtcg ggacctctgt agccgctcta tctggctagc aaggaagatt cgttcagacc    | 300  |
|      | 287 | ttgactgctc ttacggaatc ctatgtaaat tgcctatccc gctgttatct gtttccctt     | 360  |
|      | 289 | catctttttt gatccagcaa cttaccatca cgcacatcgat ccattacaa ttgtgaaagc    | 420  |
|      | 291 | tctaattcata tagtcattca tataggttat ttgacatggg cccttccctt gagaaaccc    | 480  |
|      | 293 | atgtgacttt attttcttcc tctggctgtt ttaggagatg aagttaactt aatgagaaaa    | 540  |
|      | 295 | tatatatatgaa gttctagaaaa ggatgggtt atatgtctt gaggctattt caaaatttt    | 600  |
|      | 297 | ttggccatattt attctgaata ctacctagaa cagattagcc atggccctn tgggtnttc    | 660  |
|      | 299 | ataagccatt gttctgaant ttttttagctt tgtaaatgaa aggttatgg gataffaaga    | 720  |
|      | 301 | gtnctatgaa cgtgggagga atttgtaaat cctaccaatt tntnctatattt agcattagcc  | 780  |
|      | 303 | ccccacctttt antattctgc atcaaaaatgta agatgtgtc taaagagaaaa ggttagctat | 840  |
|      | 305 | cataaggact cctataanat tcnttggaaa cttnntggaaat tttttagctt tgtaaatgaa  | 900  |
|      | 307 | tttnttggag ttccaaattt tttttagctt tgtaaatgaa aggttatgg gataffaaga     | 960  |
|      | 309 | ccccctnnng anaatgtttt gggggaaaaaa cctnccaaacc ccnttggg angaagttt     | 1020 |
|      | 311 | ttaannttt taaggcttngn ngaaacngn ttttaatttt ttgggnccnan cgcctntccc    | 1080 |
|      | 313 | cggttaccagg aaaatcagga cctnttttg ggggnngncn ccnaacngggg ggnaaaangg   | 1140 |
|      | 315 | gaaatttgcattt canaaaaat ctttccgnn nnngtgaag catcaggccc tgaacaagaa    | 1200 |
|      | 317 | catcaacctg gactctgccc atggatgcc agtggcaagc actgatcaatggatggact       | 1260 |
|      | 319 | gaccgaggca gagcgaactcc aagagaaccc tcaagctt cgtacccatcc atgtttgtt     | 1320 |
|      | 321 | ggccaggctc tttagaagacc agcagggtgca ttttacccca accgaagggtt actccatca  | 1380 |
|      | 323 | agctatacat acccttcttc tccaagtcgc tgccttgca taccagatag aggatataat     | 1440 |
|      | 325 | gatactcctg gaataacaaga tcccccgcaa tgaggctgat gggatgccta ttaatgttgg   | 1500 |

RAW SEQUENCE LISTING ERROR SUMMARY  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 39,40,41,87,650,657,679,728,762,764,792,834,858,863,874,880  
Seq#:1; N Pos. 891,904,918,927,929,935,944,954,959,967,968,969,972,994,1003  
Seq#:1; N Pos. 1012,1026,1027,1038,1040,1041,1047,1050,1066,1068,1070,1076  
Seq#:1; N Pos. 1104,1114,1115,1117,1120,1123,1126,1133,1138,1149,1153,1169  
Seq#:1; N Pos. 1170,1171,1172,1173,1174,1664,1665,1666,1667,1668  
Seq#:3; N Pos. 35,36,82,645,652,674,718,757,759,787,829,853,858,869,875,886  
Seq#:3; N Pos. 899,913,922,924,930,939,949,954,962,963,964,967,989,998,1007  
Seq#:3; N Pos. 1021,1022,1033,1035,1036,1042,1045,1061,1063,1065,1071,1099  
Seq#:3; N Pos. 1109,1110,1112,1115,1118,1121,1128,1133,1144,1148,1164,1165  
Seq#:3; N Pos. 1166,1167,1168,1169,1659,1660,1661,1662,1663  
Seq#:6; N Pos. 466,467,468,469,470,471,472,473,474,475,476,2280,2281,2282  
Seq#:6; N Pos. 2283,2284,2285,2286,2287,2288,2289,2290